

•		APR	2 6 2004	30					
	13	Ö.	, Q ²	5 /	SEQUENC	E LISTING			
	<110>	Tho Ran	Pade Ph Ph Sophi	nilip ia	Ε.				
	<120>	Sele	ected A	Antibo	ody Composi	tions for B	inding to A	minophosphol:	ipids
	<130>	400	1.00300	00					
	<140> <141>		10/621, 3-07-15						
	<150> <151>		396,263 2-07-15						
	<160>	9							
	<170>	Pate	entIn v	versio	on 3.1				
	<210> <211> <212> <213>	1 519 DNA Mus	muscul	us					
	<400>	1	cctaa	+ < + +	+2+++22+0	ctgtcagtaa	ctacaggtgt	canatatana	60
	acgggat	-gga	cccgga			ctgtcagtaa	ccacaggcgc	ccaccccyay	00
	gtccago	ctgc	agcagt	ctgg	acctgagctg	gagaagcctg	gcgcttcagt	gaagctatcc	120
	tgcaago	gctt	ctggtt	actc	attcactggc	tacaacatga	actgggtgaa	acagagccat	180
	ggaaaga	agcc	ttgaat	ggat	tggacatatt	gatccttact	atggtgatac	ttcctacaac	240
	cagaagt	tca	ggggca	aggc	cacattgact	gtagacaaat	cctccagcac	agcctacatg	300
	cagctca	aaga	gcctga	catc	tgaggactct	gcagtctatt	actgtgtaaa	ggggggttac	360
	tacgggc	cact	ggtact	tcga	tgtctggggc	gcagggacca	cggtcaccgt	ctcctcagct	420
	acaacaa	acag	ccccat	ctgt	ctatcccttg	gtcccgggcg	gatcccccgg	gctgcaggaa	480
	ttcgata	atca	agctta	tcga	taccgtcgac	ctcgagggg			519

<210> 2 <211> 152 <212> PRT

<213> Mus musculus

<400> 2

Met Gly Trp Thr Trp Ile Phe Ile Leu Ile Leu Ser Val Thr Thr Gly

Val His Ser Glu Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Glu Lys

20 25 30

Pro Gly Ala Ser Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Ser Phe 40 45 Thr Gly Tyr Asn Met Asn Trp Val Lys Gln Ser His Gly Lys Ser Leu 55 Glu Trp Ile Gly His Ile Asp Pro Tyr Tyr Gly Asp Thr Ser Tyr Asn 70 75 Gln Lys Phe Arg Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser 85 Thr Ala Tyr Met Gln Leu Lys Ser Leu Thr Ser Glu Asp Ser Ala Val 100 105 110 Tyr Tyr Cys Val Lys Gly Gly Tyr Tyr Gly His Trp Tyr Phe Asp Val 115 Trp Gly Ala Gly Thr Thr Val Thr Val Ser Ser Ala Thr Thr Ala 135 140 Pro Ser Val Tyr Pro Leu Val Pro 145 150 <210> 3 <211> 435 <212> DNA <213> Mus musculus <400> 3 atggacatga gggctcctgc acagattttg ggcttcttgt tgctcttgtt tccaggtacc 60 agatgtgaca tccagatgac ccagtctcca tcctccttat ctgcctctct gggagaaaga 120 gtcagtctca cttgtcgggc aagtcaggac attggtagta gcttaaactg gcttcagcag 180 ggaccagatg gaactattaa acgcctgatc tacgccacat ccagtttaga ttctggtgtc 240 cccaaaaggt tcagtggcag taggtctggg tcagattatt ctctcaccat cagcagcctt 300 gagtctgaag attttgtaga ctattactgt ctacaatatg ttagttctcc tcccacgttc 360 ggtgctggga ccaagctgga gctgaaacgg gctgatgctg caccaactgt cttcatcttc 420

435

gggcggatcc cccgg

```
<210> 4
<211>
       144
<212> PRT
<213> Mus musculus
<400> 4
Met Asp Met Arg Ala Pro Ala Gln Ile Leu Gly Phe Leu Leu Leu Leu
                5
Phe Pro Gly Thr Arg Cys Asp Ile Gln Met Thr Gln Ser Pro Ser Ser
                                25
Leu Ser Ala Ser Leu Gly Glu Arg Val Ser Leu Thr Cys Arg Ala Ser
                            40
Gln Asp Ile Gly Ser Ser Leu Asn Trp Leu Gln Gln Gly Pro Asp Gly
    50
                        55
Thr Ile Lys Arg Leu Ile Tyr Ala Thr Ser Ser Leu Asp Ser Gly Val
                    70
Pro Lys Arg Phe Ser Gly Ser Arg Ser Gly Ser Asp Tyr Ser Leu Thr
                85
Ile Ser Ser Leu Glu Ser Glu Asp Phe Val Asp Tyr Tyr Cys Leu Gln
            100
                                105
                                                    110
Tyr Val Ser Ser Pro Pro Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu
        115
                            120
Lys Arg Ala Asp Ala Ala Pro Thr Val Phe Ile Phe Gly Arg Ile Pro
                        135
<210> 5
<211> 783
<212> DNA
<213> ARTIFICIAL SEQUENCE
<220>
<223> SYNTHETIC OLIGONUCLEOTIDE
<400> 5
gcccagccgg ccatggccga ggtgcagctg gtggagtctg ggggaggcgt ggtccagcct
gggaggtccc tgagactctc ctgtgcagcc tctggattca ccttcagtag ctatggcatg
```

60

120

cactgggtcc	gccaggctcc	aggcaagggg	ctggagtggg	tggcagttat	atcatatgat	180
ggaagtaata	aatactatgc	agactccgtg	aagggccgat	tcaccatctc	cagagacaat	240
tccaagaaca	cgctgtatct	gcaaatgaac	agcctgagag	ctgaggacac	ggccgtgtat	300
tactgtgcaa	gattgcatgc	tcagacttgg	ggccaaggta	ccctggtcac	cgtctcgagt	360
ggtggaggcg	gttcaggcgg	aggtggctct	ggcggtagtg	cacttcagtc	tgtgctgacg	420
cagccgcctt	cagtgtctgc	ggccccagga	cagaaggtca	ccatctcctg	ctctggaagc	480
agctccgaca	tggggaatta	tgcggtatcc	tggtaccagc	agctcccagg	aacagccccc	540
aaactcctca	tctatgaaaa	taataagcga	ccctcaggga	ttcctgaccg	attctctggc	600
tccaagtctg	gcacctcagc	caccctgggc	atcactggcc	tctggcctga	ggacgaggcc	660
gattattact	gcttagcatg	ggataccagc	ccgcggaatg	tattcggcgg	agggaccaag	720
ctgaccgtcc	taggtgcggc	cgcacatcat	catcaccatc	acggggccgc	agaacaaaaa	780
ctc						783

<210> 6

<211> 261

<212> PRT

<213> ARTIFICIAL SEQUENCE

<220>

<223> POLYPEPTIDE

<400> 6

Ala Gl
n Pro Ala Met Ala Glu Val Gl
n Leu Val Glu Ser Gly Gly 1 5 10 15

Val Val Gln Pro Gly Arg Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly
20 25 30

Phe Thr Phe Ser Ser Tyr Gly Met His Trp Val Arg Gln Ala Pro Gly 35 40 45

Lys Gly Leu Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys 50 60

Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn 65 70 75 80

Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp 85 90 95

Thr Ala Val Tyr Tyr Cys Ala Arg Leu His Ala Gln Thr Trp Gly Gln
100 105 110

Gly Thr Leu Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly 115 120 125

Gly Ser Gly Gly Ser Ala Leu Gln Ser Val Leu Thr Gln Pro Pro Ser 130 135 140

Val Ser Ala Ala Pro Gly Gln Lys Val Thr Ile Ser Cys Ser Gly Ser 145 · 150 155 160

Ser Ser Asp Met Gly Asn Tyr Ala Val Ser Trp Tyr Gln Gln Leu Pro 165 170 175

Gly Thr Ala Pro Lys Leu Leu Ile Tyr Glu Asn Asn Lys Arg Pro Ser 180 185 190

Gly Ile Pro Asp Arg Phe Ser Gly Ser Lys Ser Gly Thr Ser Ala Thr
195 200 205

Leu Gly Ile Thr Gly Leu Trp Pro Glu Asp Glu Ala Asp Tyr Tyr Cys 210 215 220

Leu Ala Trp Asp Thr Ser Pro Arg Asn Val Phe Gly Gly Gly Thr Lys 225 230 235 240

Leu Thr Val Leu Gly Ala Ala Ala His His His His His Gly Ala 245 250 255

Ala Glu Gln Lys Leu 260

<210> 7

<211> 20

<212> PRT

<213> Homo sapiens

<400> 7

Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Lys 1 5 10 15

```
Ser Thr Ser Gly
            20
<210> 8
<211> 15
<212> PRT
<213> Homo sapiens
<400> 8
Ile Phe Pro Pro Ser Asp Glu Gln Leu Lys Ser Gly Thr Ala Ser
                                    10
<210> 9
<211> 19
<212> PRT
<213> Streptomyces cinnamoneus
<220>
<221> MISC FEATURE
      (11)..(18)
<222>
<223> Xaa = Abu
<400> 9
Ala Lys Gln Ala Ala Ala Phe Gly Pro Phe Xaa Phe Val Ala Asp Gly
```

Asn Xaa Lys